

AMENDMENTS TO SPECIFICATION

The additions have been indicated by underlining.

Please replace the fourth paragraph on page 10 with the following amended paragraph:

Figure 5(A, B): (a) The 3D structure of JCH2 was predicted based on homology modeling facility (Swiss-Model) available on ~~:(<http://www.expasy.org/swissmod/SWISS-MODEL.html>)~~
the Advanced Biomedical Computer Center server (NCI Frederick, USA) using the first approach mode model and (b) the deduced amino acid sequence of JCH2 [SEQ ID NO: 2] and the synthetic peptides derived from the sequence. The numbers refer to the individual peptide fragments (peptides 1-9) as shown in Table 2. Letter H, C and E refer to helix, coil and beta strand respectively.